0			2003/10/13 09:31	USPAT; US-PGPUB; EPO; JPO; DERWENT	5	9 same 6	0	L10	BRS	10
0			2003/10/13 09:31	USPAT; US-PGPUB; EPO; JPO; DERWENT	OMP	8 or hCOMP	34	L9	BRS	9
0			2003/10/13 09:31	USPAT; US-PGPUB; EPO; JPO; DERWENT		7 same human	22	L8	BRS	8
0			2003/10/13 09:31	USPAT; US-PGPUB; EPO; JPO; DERWENT	(cartilage adj oligomeric adj matrix adj protein) or (thrombospondin-5)	(cartilag protein)	52	L7	BRS	7
0			2003/10/13 09:52	USPAT; US-PGPUB; EPO; JPO; DERWENT	· · · · · · · · · · · · · · · · · · ·	elisa	40927	16	BRS	6
0			2003/10/13 09:28	USPAT; US-PGPUB; EPO; JPO; DERWENT		1 same 4	0	L5	BRS	5
0			2003/10/13 09:28	USPAT; US-PGPUB; EPO; JPO; DERWENT	("50" adj kda) or ("55" adj kda) or ("62" adj kda) or ("67" adj kda)	("50" ac adj kda)	3699	L4	BRS	4
0			2003/10/13 09:27	USPAT; US-PGPUB; EPO; JPO; DERWENT	2	1 same 2	0	L3	BRS	ω
0			2003/10/13 09:27	USPAT; US-PGPUB; EPO; JPO; DERWENT	trypsin same (cleav\$3 or digest\$3)	trypsin s	8539	L2	BRS	2
0			2003/10/13 09:30	USPAT; US-PGPUB; EPO; JPO; DERWENT	hCOMP or (cartilage adj oligomeric adj matrix adj protein) or (thrombospondin-5)	hCOMP matrix a	64	Ľ1	BRS	1
Err	Error Defin ition	Com ments	Time Stamp	DBs	Search Text		Hits	L#	Туре	

0			2003/10/13 09:53	USPAT; US-PGPUB; EPO; JPO; DERWENT		1 same 19	0	L20	BRS	20
0			2003/10/13 09:53	UB; EPO; RWENT		elisa same kit	5725	L19	BRS	19
0			2003/10/13 09:51	USPAT; US-PGPUB; EPO; JPO; DERWENT		12 same 17	2	L18	BRS	18
0			2003/10/13 09:51	USPAT; US-PGPUB; EPO; JPO; DERWENT	chondrocyte or (mesenchymal adj stem adj cell)	chondrocyte (6250	L17	BRS	17
0			2003/10/13 09:50	USPAT; US-PGPUB; EPO; JPO; DERWENT		12 same 15	0	L16	BRS	16
0			2003/10/13 09:49	USPAT; US-PGPUB; EPO; JPO; DERWENT		13 same 14	56	L15	BRS	15
0			2003/10/13 09:49	USPAT; US-PGPUB; EPO; JPO; DERWENT	(vitamin adj d3) or (retinoic adj acid)	(vitamin adj	9008	L14	BRS	14
0			2003/10/13 09:49	USPAT; US-PGPUB; EPO; JPO; DERWENT	n adj agent	differentiation adj agent	392	L13	BRS	13
0			2003/10/13 09:48	USPAT; US-PGPUB; EPO; JPO; DERWENT		1 same 11	16	L12	BRS	12
0			2003/10/13 09:33	USPAT; US-PGPUB; EPO; JPO; DERWENT	(biological adj matrix) or (treated adj cartilage) or (bone adj matrix) or collagen or hyaluronan or (fibrin adj gel) or (carbon adj fiber) or (polylactic adj acid)		123963	L11	BRS	11
Err	Error Defin ition	Com ments	Time Stamp	DBs	Search Text	and distance of the second of	Hits	L#	Туре	

	Туре	L#	Hits	Search Text	DBs	Time Stamp Com Error Err Err	Com Def	ror fin ors
21	BRS	L21	150	chen adj hui in.	USPAT; US-PGPUB; EPO; JPO; DERWENT	2003/10/13 09:54		0
22	BRS	L22	26	lawler≀adj john.in.	USPAT; US-PGPUB; EPO; JPO; DERWENT	2003/10/13 09:54		0
23	BRS	L23	-	(21 or 22) and 1	USPAT; US-PGPUB; EPO; JPO; DERWENT	2003/10/13 09:54		0

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FILE 'MEDLINE' ENTERED AT 10:01:08 N 13 OCT 2003
FILE 'CAPLUS' ENTERED AT 10:01:08 ON 13 OCT 2003
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FILE 'SCISEARCH' ENTERED AT 10:01:08 ON 13 OCT 2003
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FILE 'AGRICOLA' ENTERED AT 10:01:08 ON 13 OCT 2003
=> s hcomp or (cartilage oligomeric matrix protein) or (thrombospondin-5)
            1106 HCOMP OR (CARTILAGE OLIGOMERIC MATRIX PROTEIN) OR (THROMBOSPONDI
                  N-5
=> s trypsin (p) (cleav? or digest?)
L2
           55314 TRYPSIN (P) (CLEAV? OR DIGEST?)
=> s (50 kda) or (55 kda) or (62 kda) or (67 kda)
           35722 (50 KDA) OR (55 KDA) OR (62 KDA) OR (67 KDA)
=> s 12 (P) L3 (P) L1
L4 0 L2 (P) L3 (P) L1
=> S ELISA
         269998 ELISA
=> S L5 (P) KIT
            8257 L5 (P) KIT
=> s l1 (p) l6
               3 L1 (P) L6
=> duplicate remove 17
DUPLICATE PREFERENCE IS 'MEDLINE, BIOSIS, EMBASE'
KEEP DUPLICATES FROM MORE THAN ONE FILE? Y/(N):n
PROCESSING COMPLETED FOR L7
                1 DUPLICATE REMOVE L7 (2 DUPLICATES REMOVED)
=> d 18 1 ibib abs
     ANSWER 1 OF 1
                           MEDLINE on STN
                                                                  DUPLICATE 1
ACCESSION NUMBER:
                       2003359307
                                         MEDLINE
DOCUMENT NUMBER:
                                   PubMed ID: 12892252
TITLE:
                       Serum levels of cartilage oligomeric matrix protein. A
                       predicting factor and a valuable parameter for disease
                       management in rheumatoid arthritis.
                       Skoumal M; Kolarz G; Klingler A
Institute for Rheumatology of the Kurstadt Baden, Austria..
AUTHOR:
CORPORATE SOURCE:
                       martin.skoumal@a1.net
SOURCE:
                       SCANDINAVIAN JOURNAL OF RHEUMATOLOGY, (2003) 32 (3) 156-61.
                       Journal code: 0321213. ISSN: 0300-9742.
PUB. COUNTRY:
                       Norway
DOCUMENT TYPE:
                       (CLINICAL TRIAL)
                       Journal; Article; (JOURNAL ARTICLE)
LANGUAGE:
                       English
FILE SEGMENT:
                       Priority Journals
ENTRY MONTH:
                       200308
ENTRY DATE:
                       Entered STN: 20030802
                       Last Updated on STN: 20030809
                       Entered Medline: 20030808
                                          ***cartilage***
ΔR
     OBJECTIVE: To examine whether
                                                                  ***oligomeric***
                            ***protein*** (COMP) correlates with inflammation
        ***matrix***
     and/or joint destruction of patients with rheumatoid arthritis (RA) and to
     test COMP as predicting factor for the outcome of patients with
     established RA. METHODS: Serum levels of COMP were measured in sera of 62 patients, suffering from RA according to the ACR criteria and treated in intervals in our department, over a period of 5 years. A commercially available sandwich-type ***ELISA*** - ***kit*** developed by AnaMar
```

Medical AB, Sweden, was used. The results of serum COMP were compared with the Disease Activity Sci. (DAS), the Larsen Score, and inical and laboratory parameters. RESULTS: We found a positive correlation between serum levels of COMP at baseline and deterioration of Larsen score even after 5 years (p < 0.007; r = 0.34). To confirm serum COMP as an independent predicting factor for patients with RA we looked at a subgroup

```
of patients (n = 17) with elevated serum levels of COMP (mean 11,7 U/1)
      and low clinical prognostic factors. In this subgroup we also found a
      significant correlation with delta Larsen score (p < 0.01; r = 0.59) after
      5 years. CONCLUSION: Serum levels of COMP is known to reflect increased
      cartilage turnover. The results indicate that serum COMP may be used as a prognostic marker of cartilage degradation in a patient group with
      established RA.
=> d his
      (FILE 'HOME' ENTERED AT 10:00:47 ON 13 OCT 2003)
      FILE 'MEDLINE, CAPLUS, BIOSIS, EMBASE, SCISEARCH, AGRICOLA' ENTERED AT 10\!:\!01\!:\!08 ON 13 OCT 2003
            1106 S HCOMP OR (CARTILAGE OLIGOMERIC MATRIX PROTEIN) OR (THROMBOSPO
           55314 S TRYPSIN (P) (CLEAV? OR DIGEST?)
35722 S (50 KDA) OR (55 KDA) OR (62 KDA) OR (67 KDA)
               0 S L2 (P) L3 (P) L1
          269998 S ELISA
            8257 S L5 (P) KIT
               3 S L1 (P) L6
               1 DUPLICATE REMOVE L7 (2 DUPLICATES REMOVED)
=> s 11 (p) human
            292 L1 (P) HUMAN
=> s 19 (p) 16
              0 L9 (P) L6
=> s (biological matrix) or (treated cartilage) or (bone matrix) or collagen or hyaluronan or (fib
MISSING OPERATOR GEL) OE
The search profile that was entered contains terms or
nested terms that are not separated by a logical operator.
=> s (biological matrix) or (treated cartilage) or (bone matrix) or collagen or hyaluronan or (fib
         515579 (BIOLOGICAL MATRIX) OR (TREATED CARTILAGE) OR (BONE MATRIX) OR
                COLLAGEN OR HYALURONAN OR (FIBRIN GEL) OR (CARBON FIBER) OR
                (POLYLACTIC ACID)
=> s 11 (p) 111
            249 L1 (P) L11
=> s l12 (p) composition
             11 L12 (P) COMPOSITION
=> duplicate remove 113
DUPLICATE PREFERENCE IS 'MEDLINE, CAPLUS, BIOSIS, EMBASE, SCISEARCH'
KEEP DUPLICATES FROM MORE THAN ONE FILE? Y/(N):n
PROCESSING COMPLETED FOR L13
               4 DUPLICATE REMOVE L13 (7 DUPLICATES REMOVED)
=> d 114 1-4 ibib abs
L14 ANSWER 1 OF 4
                         MEDLINE on STN
                                                            DUPLICATE 1
                     2001011600
ACCESSION NUMBER:
                                     MEDLINE
DOCUMENT NUMBER:
                     20385047
                                PubMed ID: 10924396
TITLE:
                     Differences in the concentration of various synovial fluid
                     constituents between the distal interphalangeal joint, the
                     metacarpophalangeal joint and the navicular bursa in normal
                     horses.
AUTHOR:
                     Viitanen M; Bird J; Maisi P; Smith R; Tulamo R M; May S
CORPORATE SOURCE:
                     Farm Animal and Equine Medicine and Surgery, Royal
                     Veterinary College, University of London, UK.
                     RESEARCH IN VETERINARY SCIENCE, (2000 Aug) 69 (1) 63-7.
SOURCE:
                     Journal code: 0401300. ISSN: 0034-5288.
PUB. COUNTRY:
                     ENGLAND: United Kingdom
DOCUMENT TYPE:
                     Journal; Article; (JOURNAL ARTICLE)
LANGUAGE:
                     English
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Priority Journals FILE SEGMENT: ENTRY MONTH: 200010

L1

L2 L3 L4

L5

L6 L7

--L8--

L10

L11

L12

ENTRY DATE:

Entered STN: 20010322 Last Updated STN: 20010322 Entered Medline: 20001023

As a prerequisite for the identification of navicular disease markers, the concentrations of ***cartilage*** ***oligomeric*** ***matrix*** AB ***protein*** (COMP), total glycosaminoglycans (GAG), ***hyalurometalloproteinases (MMP) 2 and 9 and total protein were measured in ***hyaluronan*** synovial fluid samples obtained from the distal interphalangeal joint (DIP), the metacarpophalangeal joint (MCP) and the navicular bursa of 24 horses. Mean GAG, COMP and total protein levels were significantly higher in the DIP joint and in the navicular bursa compared to the MCP joint. ***Hyaluronan*** content was lower. MMP -2 activity was present in all fluids measured and had similar levels in different joints. MMP -9 was present in 42 per cent of MCP joint samples and 58 per cent of DIP joint samples and of navicular bursal samples. In relation to the constituents measured, the ***composition*** of navicular bursal fluid was similar to the articular synovial fluids, in particular that obtained from the DIP joint. Correlation between the constituents of DIP joint fluid and navicular bursal fluid obtained from the same legs was statistically significant for all the parameters measured.

L14 ANSWER 2 OF 4 MEDLINE on STN 2000124477 ACCESSION NUMBER: MEDLINE

DOCUMENT NUMBER: 20124477 PubMed ID: 10659252

TITLE:

Should equine athletes commence training during skeletal development?: changes in tendon matrix associated with development, ageing, function and exercise.

Smith_R_K;_Birch_H;_Patterson-Kane_J; Firth_E_C; Williams AUTHOR:

CORPORATE SOURCE:

L; Cherdchutham W; van Weeren W R; Goodship A E
Royal Veterinary College, Hatfield, Herts, UK.
EQUINE VETERINARY JOURNAL. SUPPLEMENT, (1999 Jul) 30 201-9.
Journal code: 9614088. SOURCE:

ENGLAND: United Kingdom

Journal; Article; (JOURNAL ARTICLE) DOCUMENT TYPE:

LANGUAGE: English

PUB. COUNTRY:

FILE SEGMENT: Priority Journals

ENTRY MONTH: 200003

ENTRY DATE: Entered STN: 20000314

Last Updated on STN: 20000314 Entered Medline: 20000302

In human athletes, conditioning, training and competition are commenced before skeletal maturity. Yet in equine athletics, racing of young (age 2 years) horses remains contentious. Tendon injury persists as major causes of wastage in equine athletes. Minimising injury and associated welfare AB issues could involve a radical approach to the timing and implementation of conditioning and training. Tendons were examined from Thoroughbreds, Dutch Warmblood foals, working horses and also a group of wild horses to evaluate effects of age, function and exercise. Gross mechanical properties did not differ significantly with age or exercise, but showed a high variance within each group. Mechanical properties of tendon tissue showed significant differences as a function of age and location. The ***collagen*** fibril crimp angle and length showed a regional reduction in the central core with exercise and age, with a some officers.

in the central core with exercise and age, with a synergistic effect. Regional differences in ***collagen*** fibril diameter were seen fibril diameter were seen in long-term exercised older horses, but not in short-term exercised, or younger, horses. The higher proportion of small fibrils in the central region of the long-term exercised horses did not correlate with new ***collagen*** formation and therefore appear to result from disassembly of the larger diameter fibrils. Fibril diameter distributions were

influenced by exercise regimens in the growing foal. Changes in molecular

composition occurred in longer-term exercise and older horses, in
the centre of the tendon, with higher levels of type III

collagen

the centre of the tendon, with higher levels of type III ***coll and changes in glycosaminoglycan (GAG) content. ***Cartilage*** and changes in glycosaminoglycan (GAG) content.

Oligomeric

Matrix

Prote ***Protein (COMP) levels also appear to be modulated by age, function and superimposition of exercise. These changes were all exacerbated with age and exercise, suggesting appropriate exercise in young horses may lead to a lower incidence of injury than in older horses. An hypothesis is advanced that immature tendon can respond to exercise while mature tendon has limited, if any, ability to do so. These findings support potentially controversial

DUPLICATE 2

earlier conditioning and racing of younger, rather than older, equine athletes.

L14 ANSWER 3 OF 4 ACCESSION NUMBER:

MEDLINE on STN 96195288 MEDLINE

96195288 PubMed ID: 8619919

DOCUMENT NUMBER: Predictors of joint damage in rheumatoid arthritis. TITLE:

AUTHOR: Wollheim F A Department of eumatology, Lund University Ho CORPORATE SOURCE: tal,

Sweden.

APMIS, (1996 Feb) 104 (2) 81-93. Ref: 103 Journal code: 8803400. ISSN: 0903-4641. SOURCE:

PUB. COUNTRY: Denmark

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

General Review; (REVIEW)

(REVIEW, ACADEMIC)

LANGUAGE: English

Priority Journals FILE SEGMENT:

ENTRY MONTH: 199606

ENTRY DATE:

Entered STN: 19960627 Last Updated on STN: 19980206

Entered Medline: 19960614

AB Rheumatoid arthritis (RA) is the dominant form of destructive chronic arthritis with the potential to cause substantial disability and permanent functional impairment. The final extent and progression rate with time, however, varies markedly. In order to study effects of intervention and to support early aggressive and atoxic therapy in selected cases, predictive disease markers are needed. Recent advances regarding joint tissue ***composition*** and pathophysiology have defined a number of tissue ***composition*** and pathophysiology have defined a number of biological marker candidates which need to be explored for possible prognostic information. Some markers are characteristic for RA, such as rheumatoid factors and certain autoantibodies, which although they are more prevalent among patients with aggressive disease are not sensitive as predictors in early disease. Genetic susceptibility markers have been claimed to be good predictors of persisting arthritis in early synovitis clinics, but their role as severity markers in established disease is limited. Unspecific markers of inflammation, notably ESR or CRP when persistently elevated, are useful to monitor disease course and newer markers need to document their superiority over these. Another group of markers are attractive because of enriched or exclusive occurrence in joint tissue, and altered metabolism in joint disease. Thus, ***collagen*** type III propeptides, hyaluronates, and neopterin originating in the synovium could be useful, and, in particular, hyaluronate levels indeed do provide some predictive information. Highly tissue-specific cartilage metabolites include aggrecan fragments, ***collagen*** II fragments, ***cartilage*** ***oligomeric*** ***matrix*** ***protein*** (COMP) and the extraarticular cartilage matrix protein (CMP). When used alone or in combination in early disease some information can be obtained which may in the future facilitate

prognostication. Bone metabolism can be monitored and there are different markers for synthesis and resorption. Meanwhile, whilst the new markers are essential research tools, their routine clinical usefulness remains to be proven.

L14 ANSWER 4 OF 4 MEDLINE on STN DUPLICATE 3

ACCESSION NUMBER: 93079835 MEDLINE

DOCUMENT NUMBER: 93079835 PubMed ID: 1448898

Immunohistochemical localization of matrix proteins in the TITLE:

femoral joint cartilage of growing commercial pigs.

AUTHOR: Ekman S; Heinegard D

CORPORATE SOURCE: Department of Anatomy and Histology, Swedish University of

Agricultural Sciences, Uppsala.

VETERINARY PATHOLOGY, (1992 Nov) 29 (6) 514-20. Journal code: 0312020. ISSN: 0300-9858. SOURCE:

PUB. COUNTRY: United States

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 199212

ENTRY DATE: Entered STN: 19930129

Last Updated on STN: 19930129 Entered Medline: 19921228

The immunocytochemical localization of several matrix macromolecules ΔR including ***collagen*** type II and proteoglycans, in the distal femoral articular-epiphyseal cartilage complex of 15 commercial pigs between the age of 6 and 18 weeks was studied. Early osteochondrotic lesions, i.e., chondronecrosis in the resting region of the growth cartilage, as well as extensions of necrotic cartilage into the subchondral bone, were present in all animals, except those 6 weeks old. A battery of antibodies were used for identification of macromolecules in the matrix at different stages of the disease. Chondrocyte involvement in the process could be studied by identifying the sequence of alterations in matrix macromolecules as the lesion developed. The immunostaining for aggrecan (large aggregating proteoglycans), ***cartilage***

```
***oligomeric*** ***matrix*** ***protein*** , fibronectin,
***collagen*** type II, fromodulin, and biglycan was more prominent in
the areas of chondronecrosis, extending into the subchondral bone, than in
       the normal resting region. This altered pattern of matrix macromolecules resembled that of the matrix of the proliferative chondrocytes and
       suggests that the chondrocyte maturation had stopped in the proliferative
       zone. The matrix in the areas of chondronecrosis in the resting region resembled that in the normal resting region. Thus the chondronecrosis
       appears to have preceded alterations of the matrix
                                                                              ***composition***
       The antibody reactivity pattern was, however, altered in the matrix of the
       clustered chondrocytes in areas of chondronecrosis. Staining in these
       regions suggested a more prominent appearance of fibronectin and
       ***collagen*** type II than in the normal matrix of the resting region. These changes are suggestive of attempt to repair.(ABSTRACT TRUNCATED AT
       250 WORDS)
=> d his
       (FILE 'HOME' ENTERED AT 10:00:47 ON 13 OCT 2003)
       FILE 'MEDLINE, CAPLUS, BIOSIS, EMBASE, SCISEARCH, AGRICOLA' ENTERED AT 10:01:08 ON 13 OCT 2003

1106 S HCOMP OR (CARTILAGE OLIGOMERIC MATRIX PROTEIN) OR (THROMBOSPO
             55314 S TRYPSIN (P) (CLEAV? OR DIGEST?)
-35722 S (50 KDA) OR (55 KDA) OR (62 KDA) OR (67 KDA)
0 S L2 (P) L3 (P) L1
            269998 S ELISA
               8257 S L5 (P) KIT
                   3 S L1 (P) L6
                   1 DUPLICATE REMOVE L7 (2 DUPLICATES REMOVED)
                292 S L1 (P) HUMAN
                   0 S L9 (P) L6
            515579 S (BIOLOGICAL MATRIX) OR (TREATED CARTILAGE) OR (BONE MATRIX) O
                249 S L1 (P) L11
                  11 S L12 (P) COMPOSITION
                   4 DUPLICATE REMOVE L13 (7 DUPLICATES REMOVED)
=> s 112 (p) (purified)
                10 L12 (P) (PURIFIED)
=> duplicate remove 115
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PROCESSING COMPLETED FOR L15
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=> s 116 not 114
                 2 L16 NOT L14
=> d 117 1-2 ibib abs
L17 ANSWER 1 OF 2
                               MEDLINE on STN
ACCESSION NUMBER:
                           2003324085
                                               IN-PROCESS
DOCUMENT NUMBER:
                           22737940
                                         PubMed ID: 12853037
TITLE:
                           Cleavage of cartilage oligomeric matrix protein
                           (thrombospondin-5) by matrix metalloproteinases and a disintegrin and metalloproteinase with thrombospondin
                          motifs.
AUTHOR:
                          Dickinson Sally C; Vankemmelbeke Mireille N; Buttle David
                           J; Rosenberg Krisztina; Heinegard Dick; Hollander Anthony P
                          Academic Rheumatology, University of Bristol, Avon
CORPORATE SOURCE:
                          Orthopaedic Centre, Southmead Hospital, BS10 5NB, Bristol,
SOURCE:
                          MATRIX_BIOLOGY, (2003 May) 22 (3) 267-78.
                           Journal code: 9432592. ISSN: 0945-053x.
PUB. COUNTRY:
                          Germany: Germany, Federal Republic of Journal; Article; (JOURNAL ARTICLE)
DOCUMENT TYPE:
LANGUAGE:
                          English
FILE SEGMENT:
                          IN-PROCESS; NONINDEXED; Priority Journals
                          Entered STN: 20030711
ENTRY DATE:
                          Last Updated on STN: 20030808
         ***Cartilage***
                                    ***oligomeric***
                                                                ***matrix***
                                                                                       ***protein***
      (COMP) is a pentameric glycoprotein present in cartilage, tendon and ligament. Fragments of the molecule are present in the diseased cartilage, synovial fluid and serum of patients with knee injuries,
```

osteoarthritis and rheumatoid arthritis. Although COMP is a substrate for

L1

L2 -Ŀ3· L4 L5

L6 L7

L8 L9

L10

L11 L12

L13

L14

L15

several matrix metalloproteinses (MMPs), the enzymes responsible for COMP degradation in vivo have yet be identified. In this study well-established bovine cartilage culture models to examine IL-lalpha-stimulated COMP proteolysis in the presence and absence of MMP inhibitors. COMP was released from bovine nasal cartilage, in response to IL-lalpha, at an intermediate time between proteoglycans and type II ***collagen***, when soluble MMP levels in the culture medium were undetectable. The major fragment of COMP released following IL-1alpha-stimulation migrated with an apparent molecular mass of approximately 110 kDa (Fragment-110) and co-migrated with both the major fragment present in human arthritic synovial fluid samples and the product of COMP cleavage by ***purified*** MMP-9. However, the broad-spectrum MMP-9. However, the broad-spectrum MMP and ADAM inhibitor BB94 only partially inhibited the formation of Fragment-110 and failed to inhibit COMP release significantly. Therefore the results of these studies indicate a role for proteinases other than MMPs in the degradation of COMP in bovine cartilage. It was further demonstrated that ***purified*** COMP was cleaved by ADAMTS-4, but not ADAMTS-1 or -5, to yield a fragment which co-migrated with Fragment-110. Therefore this is the first demonstration of COMP as a substrate for ADAMTS-4, although it remains to be determined whether this enzyme plays a role in COMP degradation in vivo.

L17 ANSWER 2 OF 2 MEDLINE on STN ACCESSION NUMBER: 1998378148 MEDLINE

DOCUMENT NUMBER: 98378148 PubMed ID: 9714346

Analysis of cartilage oligomeric matrix protein (COMP) in TITLE:

synovial fibroblasts and synovial fluids.

Hummel K_M; Neidhart_M; Vilim V; Hauser N; Aicher W K; Gay AUTHOR:

R E; Gay S; Hauselmann H J Center for Experimental Rheumatology, University Hospital, CORPORATE SOURCE:

Zurich, Switzerland.

BRITISH JOURNAL OF RHEUMATOLOGY, (1998 Jul) 37 (7) 721-8. SOURCE:

Journal code: 8302415. ISSN: 0263-7103.

PUB. COUNTRY: ENGLAND: United Kingdom

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

Abridged Index Medicus Journals; Priority Journals FILE SEGMENT:

ENTRY MONTH: 199809

ENTRY DATE:

Entered STN: 19980917 Last Updated on STN: 19980917 Entered Medline: 19980904

We investigated the expression of ***matrix*** ***protein*** ***cartilage*** AB ***oligomeric*** ***matrix*** ***protein*** (COMP) in normal and rheumatoid arthritis (RA) synovial fibroblasts. In situ hybridization (ISH) was conducted on synovial specimens from five RA patients applying specific probes for COMP ***collagen*** or fibroblast type I. ISH was combined with immunohistochemistry, applying antibodies to the macrophage marker CD68. Ribonuclease protection assay (RPA) and rapid amplification of 3'-cDNA ends (3'-RACE) were performed on total RNA from normal and RA synovial fibroblast cultures. Protein extracts from fibroblasts and culture supernatants were compared with synovial fluids and protein extracts from isolated chondrocytes by Western blot utilizing polyclonal and monoclonal antibodies (18-G3 mAb) to COMP. COMP mRNA was detected in fibroblasts of RA synovium by ISH, and in normal and RA synovial fibroblast cultures by RPA. 3'-RACE demonstrated sequence homology of chondrocyte and synovial fibroblast COMP along the coding sequence. COMP protein was detected in synovial fibroblasts and culture supernatants by immunoblot. Using polyclonal antibodies, the major portion of COMP from fibroblasts and culture supernatants was present as low-molecular-weight (LMW) bands, corresponding to those found in synovial fluids. These LMW COMP bands, however, were not detected in any of the cells or tissues tested using -G3 mAb. In protein extracts from chondrocytes and in COMP
purified from cartilage, these LMW bands could not be detected. In conclusion, the data suggest that certain forms of COMP detected in synovial fluid are secreted from synovial fibroblasts and could be

distinguished by specific mAbs from COMP secreted by chondrocytes.

=> d his

(FILE 'HOME' ENTERED AT 10:00:47 ON 13 OCT 2003)

FILE 'MEDLINE, CAPLUS, BIOSIS, EMBASE, SCISEARCH, AGRICOLA' ENTERED AT 10:01:08 ON 13 OCT 2003

1106 S HCOMP OR (CARTILAGE OLIGOMERIC MATRIX PROTEIN) OR (THROMBOSPO 55314 S TRYPSIN (P) (CLEAV? OR DIGEST?) 35722 S (50 KDA) OR (55 KDA) OR (62 KDA) OR (67 KDA)

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0 S L2 (P) L3 (P) L1
            269998 S ELISA
 L5
 L6
              8257 S L5 (P) KIT
                    S L1 (P) L6
                  1 DUPLICATE REMOVE L7 (2 DUPLICATES REMOVED)
 L9
                292 S L1 (P) HUMAN
 L10
                  0 S L9 (P) L6
            515579 S (BIOLOGICAL MATRIX) OR (TREATED CARTILAGE) OR (BONE MATRIX) O
 L11
                249 S L1 (P) L11
                 11 S L12 (P) COMPOSITION
 L13
                  4 DUPLICATE REMOVE L13 (7 DUPLICATES REMOVED)
 L14
                 10 S L12 (P) (PURIFIED)
 L15
                  2 DUPLICATE REMOVE L15 (8 DUPLICATES REMOVED)
 L16
                  2 S L16 NOT L14
 L17
 => s chondrocyte or (mesenchymal stem cell)
 L18
            51010 CHONDROCYTE OR (MESENCHYMAL STEM CELL)
 => s differentiation agent
 L 19
              831 DIFFERENTIATION AGENT
 => s (vitamin d) or (retinoic acid)
 L20
          210357 (VITAMIN D) OR (RETINOIC ACID)
 => duplicate remove 112
-DUPLICATE PREFERENCE IS MEDLINE, CAPLUS, BIOSIS, EMBASE, SCISEARCH, AGRICOLA'
 KEEP DUPLICATES FROM MORE THAN ONE FILE? Y/(N):n
 PROCESSING COMPLETED FOR L12
                78 DUPLICATE REMOVE L12 (171 DUPLICATES REMOVED)
 => s 121 (p) 118
 PROXIMITY OPERATOR LEVEL NOT CONSISTENT WITH
 FIELD CODE - 'AND' OPERATOR ASSUMED 'L142 (P) L117'
 PROXIMITY OPERATOR LEVEL NOT CONSISTENT WITH
 FIELD CODE - 'AND' OPERATOR ASSUMED 'L146 (P) L119'
               14 L21 (P) L18
 L22
=> s 122 not (117 or 114)
 L23
               12 L22 NOT (L17 OR L14)
=> d 123 1-12 ibib abs
L23 ANSWER 1 OF 12
                              MEDLINE on STN
ACCESSION NUMBER:
                         2003346330
                                            MEDLINE
DOCUMENT NUMBER:
                         22760698
                                      PubMed ID: 12878157
TITLE:
                         Redifferentiation of dedifferentiated chondrocytes and
                         chondrogenesis of human bone marrow stromal cells via
                         chondrosphere formation with expression profiling by
                         large-scale cDNA analysis.
Imabayashi Hideaki; Mori Taisuke; Gojo Satoshi; Kiyono
Tohru; Sugiyama Tomoyasu; Irie Ryotaro; Isogai Takao; Hata
Jun-ichi; Toyama Yoshiaki; Umezawa Akihiro
AUTHOR:
CORPORATE SOURCE:
                         National Research Institute for Child Health and
                         Development, Tokyo 157-8535, Japan.
EXPERIMENTAL CELL RESEARCH, (2003 Aug 1) 288 (1) 35-50.
Journal code: 0373226. ISSN: 0014-4827.
SOURCE:
                         United_States
PUB. COUNTRY:
DOCUMENT TYPE:
                         Journal; Article; (JOURNAL ARTICLE)
LANGUAGE:
                         English
FILE SEGMENT:
                         Priority Journals
ENTRY MONTH:
                         200309
ENTRY DATE:
                         Entered STN: 20030725
                         Last Updated on STN: 20030925
                         Entered Medline: 20030924
ΑB
      Characterization of dedifferentiated
                                                      ***chondrocytes***
                                                                                (DECs) and
                                   ***stem***
         ***mesenchymal***
                                                      ***cells***
                                                                      capable of
                                  ***chondrocytes*** is of biological and clinical
      differentiating into
      interest. We isolated DECs and bone marrow stromal cells (BMSCs), H4-1
      and H3-4, and demonstrated that the cells started to produce extracellular matrices, such as type II ***collagen*** and aggrecan, at an early
      matrices, such as type II ***collagen*** and aggrecan, at an early stage of chondrosphere formation. Furthermore, cDNA sequencing of cDNA
       libraries constricted by the oligocapping method was performed to analyze
      difference in mRNA expression profiling between DECs and marrow stromal cells. Upon redifferentiation of DECs, cartilage-related extracellular matrix genes, such as those encoding leucine-rich small proteoglycans, ***cartilage*** ***oligomeric*** ***matrix*** ***protein*
                                                                                ***protein***
      and chitinase 3-like 1 (cartilage glycoprotein-39), were highly expressed.
```

Growth factors such as FGF7 and CTGF were detected at a high frequency in the growth stage of monolaye tromal cultures. By combining expression profile and flow cytometry, we demonstrated that isolated stromal cells, defined by CD34(-), c-kit(-), and CD140alpha(- or low), have chondrogenic potential. The newly established human mesenchymal cells with expression profiling provide a powerful model for a study of chondrogenic differentiation and further understanding of cartilage regeneration in the means of redifferentiated DECs and BMSCs.

L23 ANSWER 2 OF 12 MEDLINE on STN

2003243202 ACCESSION NUMBER: **IN-PROCESS**

DOCUMENT NUMBER: 22650296 PubMed ID: 12766479

Apoptosis staining in cultured pseudoachondroplasia TITLE:

chondrocytes.

Duke J; Montufar-Solis D; Underwood S; Lalani Z; Hecht J T **AUTHOR:** CORPORATE SOURCE: Department of Orthodontics, Dental Branch, The University

of Texas Health Science Center at Houston..

Pauline.J.Duke@uth.tmc.edu

SOURCE: APOPTOSIS, (2003 Mar) 8 (2) 191-7.

Journal code: 9712129. ISSN: 1360-8185.

United States PUB. COUNTRY:

Journal; Article; (JOURNAL ARTICLE) DOCUMENT TYPE:

LANGUAGE: English

FILE SEGMENT: IN-PROCESS; NONINDEXED; Priority Journals

Entered STN: 20030528 ENTRY DATE:

Last Updated on STN: 20030528

Pseudoachondroplasia (PSACH) is a skeletal dysplasia caused by a mutation in ***cartilage*** ***oligomeric*** ***matrix***

cartilage ***oligomeric*** ***matrix***

protein (COMP), a glycoprotein of normal cartilage matrix.

chondrocytes have a distinctive phenotype with enlarged ren cisternae containing COMP, aggrecan, type IX ***collagen***, and chaperone proteins. Ultrastructural studies suggested that this accumulation compromises cell function, hastening cell death, and consequently reducing the number of cells in the growth plate contributing to linear bone growth. Using the alginate bead system, we cultured control and PSACH ***chondrocytes*** for twenty weeks and one year to determine the effect of the mutation on size and number of cartilage nodules; and the presence of apoptotic cell death (TUNEL assay). weeks, beads containing PSACH or control ***chondrocytes*** d weeks, beads containing PSACH or control ***chondrocytes*** did not differ in size and number of cartilage nodules or number of TUNEL-positive cells. After one year, nodule number, size and percent cartilage per bead were significantly less in PSACH nodules, and the number of cells staining positive for apoptosis was significantly greater number of cells (71.8%) vs. 44.6%). The increase in apoptosis in PSACH nodules correlates with a decrease in growth of cartilage, supporting our hypothesis that death of damaged cells contributes to the growth plate defects in PSACH.

ANSWER 3 OF 12 MEDLINE on STN

ACCESSION NUMBER: 2002432171 MEDLINE

DOCUMENT NUMBER: 22176769 PubMed ID: 12189245

Pseudoachondroplasia is caused through both intra- and TITLE:

extracellular pathogenic pathways.

Dinser Robert; Zaucke Frank; Kreppel Florian; Hultenby **AUTHOR:**

Kjell; Kochanek Stefan; Paulsson Mats; Maurer Patrik Institute for Biochemistry II, University of Cologne,

Cologne, Germany.. robert.dinser@uniklinik-saarland.de SOURCE:

JOURNAL OF CLINICAL INVESTIGATION, (2002 Aug) 110 (4)

505-13.

Journal code: 7802877. ISSN: 0021-9738.

PUB. COUNTRY: United States

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English FILE SEGMENT: ENTRY MONTH:

CORPORATE SOURCE:

ΑB

Abridged Index Medicus Journals; Priority Journals

200209

ENTRY DATE:

Entered STN: 20020822 Last Updated on STN: 20020907 Entered Medline: 20020906

Pseudoachondroplasia is a dominantly inherited chondrodysplasia associated with mutations in ***cartilage*** ***oligomeric*** ***matrix*** ***oligomeric*** ***matrix***

with mutations in ***protein*** (COMP). Investigations into the pathogenesis of pseudoachondroplasia are hampered by its rarity. We developed a cell

culture model by expressing mutant COMP in bovine primary

chondrocytes using a gutless adenoviral vector. We show
mutant COMP exerts its deleterious effects through both intra- and We show that extracellular pathogenic pathways. Overexpression of mutant COMP led to a dose-dependent decrease in cellular viability. The secretion of mutant COMP was markedly delayed, presumably due to a prolonged association with

chaperones in the endoplasmic reticulum (ER). The ECM lacked organized ***collagen*** fibers and howed amorphous aggregates for by mutant COMP. Thus, pseudoachondroplasia appears to be an ER storage disease, most likely caused by improper foliand by mutant COMP. The growth failure of affected patients may be explained by an increased cell death of growth-plate ***chondrocytes***. Dominant interference of the mutant protein on ***collagen*** fiber assembly could contribute to the observed failure of the ECM of cartilage and tendons.

L23 ANSWER 4 OF 12 MEDLINE on STN 2002092991 **ACCESSION NUMBER:** MEDLINE

DOCUMENT NUMBER: 21656885 PubMed ID: 11798989

Autologous chondrocyte transplantation. Biomechanics and TITLE:

long-term durability.

Peterson Lars; Brittberg Mats; Kiviranta Illka; Akerlund **AUTHOR:**

Evy Lundgren; Lindahl Anders

CORPORATE SOURCE: Gothenburg Medical Center, Gothenburg University,

Gothenburg, Sweden.

AMERICAN JOURNAL OF SPORTS MEDICINE, (2002 Jan-Feb) 30 (1) SOURCE:

2-12.

Journal code: 7609541. ISSN: 0363-5465.

PUB. COUNTRY: **United States**

(EVALUATION STUDIES) DOCUMENT TYPE:

Journal; Article; (JOURNAL ARTICLE)

English LANGUAGE:

-FILE-SEGMENT: Priority Journals

ENTRY MONTH: 200203

ENTRY DATE: Entered STN: 20020202

Last Updated on STN: 20020302 Entered Medline: 20020301

We evaluated the durability of autologous ***chondrocyte***
transplantation grafts in 61 patients treated for isolated cartilage
defects on the femoral condyle or the patella and followed up for a mean
of 7.4 years (range, 5 to 11). Durability was determined by comparing the
clinical status at the long-term follow-up with that found 2 years after
the transplantation. After 2 years, 50 of the 61 patients had good or
excellent clinical results, and 51 of 61 had good or excellent results at
5 to 11 years later. Grafted areas from 11 of the patients were evaluated
with an electromechanical indentation probe during a second-look with an electromechanical indentation probe during a second-look arthroscopy procedure (mean follow-up, 54.3 months; range, 33 to 84); stiffness measurements were 90% or more of those of normal cartilage in eight patients. Eight of twelve 2-mm biopsy samples taken from these patients showed hyaline characteristics with safranin O staining and a homogeneous appearance in polarized light. The constant to hyaline biopsy specimens stained positive to aggrecan and to hyaline biopsy specimens stained positive to aggrecan and to hyaline biopsy specimens stained positive to aggrecan and to hyaline biopsy specimens are highly accountable to the homogeneous appearance in polarized light. Hyaline-like specimens stained positive for type II ***collagen*** and fibrous, for type I ***collagen*** . Autologous ***chondrocyte*** transplantation for the treatment of articular

cartilage injuries has a durable outcome for as long as 11 years.

L23 ANSWER 5 OF 12 MEDLINE on STN 2002026731 ACCESSION NUMBER: MEDLINE

DOCUMENT NUMBER: 21363816 PubMed ID: 11470401

TITLE: Calreticulin, PDI, Grp94 and BiP chaperone proteins are

associated with retained COMP in pseudoachondroplasia

chondrocytes.

Hecht J T; Hayes E; Snuggs M; Decker G; Montufar-Solis D; Doege K; Mwalle F; Poole R; Stevens J; Duke P J University of Texas Medical School at Houston, Department **AUTHOR:**

CORPORATE SOURCE:

of Pediatrics, P.O. Box 20708, Houston, TX 77225-0708,

USA.. jacqueline.t.hecht@uth.tmc.edu

MATRIX BIOLOGY, (2001 Jul) 20 (4) 251-62. Journal code: 9432592. ISSN: 0945-053x. SOURCE:

Germany: Germany, Federal Republic of Journal; Article; (JOURNAL ARTICLE) PUB. COUNTRY: DOCUMENT TYPE:

English

Priority Journals; Space Life Sciences

ENTRY MONTH: 200112

LANGUAGE:

AB

FILE SEGMENT:

ENTRY DATE: Entered STN: 20020121

Last Updated on STN: 20020131

Entered Medline: 20011207
** ***oligomeric*** ***Cartilage*** ***matrix*** ***protein*** (COMP), a large pentameric glycoprotein and member of the thrombospondin (TSP) group of extracellular proteins, is found in the territorial matrix surrounding ***chondrocytes*** . More than 50 unique COMP mutations have been identified as causing two skeletal dysplasias:

pseudoachondroplasia (PSACH); and multiple epiphyseal dysplasia (EDM1). Recent studies suggest that cium-binding and calcium-induce protein folding differ between wild type and mutant proteins, and abnormal processing of the mutant COMP protein contributes to the characteristic enlarged lamellar appearing TER cisternae in PSACH and EDMI

chondrocytes in vivo and in vitro. Towards the goal of delineating the nathogenesis of BSACH and EDM1 in vivo BSACH arguments.

delineating the pathogenesis of PSACH and EDM1, in-vivo PSACH growth plate and in-vitro PSACH ***chondrocytes*** cultured in alginate beads were examined to identify and localize the chaperone proteins participating in the processing of the retained extracellular matrix proteins in the PSACH

rER. Aggrecan was localized to both the rER cisternae and matrix while COMP and type IX ***collagen*** were only found in the rER. Type II ***collagen*** was solely found in the ECM suggesting that it is processed and transported differently from other retained ECM proteins. Five chaperone proteins: BiP (Grp78); calleticulin (CRT); protein disulfide (PDI); ERp72; and Grp94, demonstrated immunoreactivity in the

enlarged PSACH cisternae and the short rER channels of

*** chondrocytes *** from both in-vivo and in-vitro samples. The chaperone proteins cluster around the electron dense material within the enlarged rER cisternae. CRT, PDI and GRP94 AB-gold particles appear to be closely associated with COMP. Immunoprecipitation and Western blot, and Fluorescence Resonance Energy Transfer (FRET) analyses indicate that CRT, PDI and GRP94 are in close proximity to normal and mutant COMP and BiP to mutant COMP. These results suggest that these proteins play a role in the processing and transport of wild type COMP in normal ***chondrocytes*** ***chondrocytes*** and in the retention of mutant COMP in PSACH

ANSWER 6 OF 12 MEDLINE on STN ACCESSION NUMBER: 2001640896 MEDLINE

DOCUMENT NUMBER: 21550102 PubMed ID: 11691584

Selective intracellular retention of extracellular matrix TITLE:

proteins and chaperones associated with

pseudoachondroplasia. **AUTHOR:**

Vranka J; Mokashi A; Keene D R; Tufa S; Corson G; Sussman

M; Horton W A; Maddox K; Sakai L; Bachinger H P CORPORATE SOURCE: Research Department, Shriners Hospital for Children,

Portland, OR 97201, USA. AR45582 (NIAMS)

CONTRACT NUMBER:

MATRIX BIOLOGY, (2001 Nov) 20 (7) 439-50. Journal code: 9432592. ISSN: 0945-053x. SOURCE:

Germany: Germany, Federal Republic of Journal; Article; (JOURNAL ARTICLE) PUB. COUNTRY: DOCUMENT TYPE:

LANGUAGE: English

FILE SEGMENT: ENTRY MONTH: Priority Journals

200202 ENTRY DATE:

Entered STN: 20011107 Last Updated on STN: 20020205 Entered Medline: 20020204

cartilage Mutations in the ***oligomeric*** ***matrix*** ***protein*** (COMP) gene result in pseudoachondroplasia (PSACH), which is a chondrodysplasia characterized by early-onset osteoarthritis and short stature. COMP is a secreted pentameric glycoprotein that belongs to the thrombospondin family of proteins. We have identified a novel missense mutation which substitutes a glycine for an aspartic acid residue in the thrombospondin (TSP) type 3 calcium-binding domain of COMP in a patient diagnosed with PSACH. Immunohistochemistry and immunoelectron microscopy both show abnormal retention of COMP within characteristically enlarged rER inclusions of PSACH. ***chondrocytes***, as well as retention of fibromodulin, decorin and types IX, XI and XII
collagen Aggregation and types IX.

collagen Aggrecan and types II and VI ***collagen*** not retained intracellularly within the same cells. In addition to selective extracellular matrix components, the chaperones HSP47, protein disulfide isomerase (PDI) and calnexin were localized at elevated levels within the rER vesicles of PSACH ***chondrocytes***, suggesting that they may play a role in the cellular retention of mutant COMP molecules. Whether the aberrant rER inclusions in PSACH ***chondrocytes*** are direct consequence of chaperone-mediated retention of mutant COMP or are otherwise due to selective intracellular protein interactions, which may in turn lead to aggregation within the rER, is unclear. However, our data demonstrate that retention of mutant COMP molecules results in the selective retention of ECM molecules and molecular chaperones, indicating the existence of distinct secretory pathways or ER-sorting mechanisms for matrix molecules, a process mediated by their association with various molecular chaperones.

ANSWER 7 OF 12 MEDLINE on STN ACCESSION NUMBER: 2001439865 MEDLINE

378166 PubMed ID: 11485547
***Cartilag
protein (COMP) and DOCUMENT NUMBER: 21378166 ***oligomeric*** rix*** TITLE: ***collagen*** īx are sensitive markers for the differentiation state of articular primary ***chondrocytes*** . articular primary ***chondrocytes***
Zaucke F; Dinser R; Maurer P; Paulsson M
Institute for Biochemistry II, Medical Faculty, University **AUTHOR:** CORPORATE SOURCE: of Cologne, Joseph-Stelzmann-Strasse 52, D-50931 Cologne. Germany.. frank.zaucke@uni-koeln.de BIOCHEMICAL JOURNAL, (2001 Aug 15) 358 (Pt 1) 17-24. Journal code: 2984726R. ISSN: 0264-6021. SOURCE: England: United Kingdom
Journal; Article; (JOURNAL ARTICLE) PUB. COUNTRY: DOCUMENT TYPE: LANGUAGE: English FILE SEGMENT: Priority Journals ENTRY MONTH: 200109 **ENTRY DATE:** Entered STN: 20010924 Last Updated on STN: 20010924 Entered Medline: 20010920 Primary ***chondrocytes*** dedifferentiate in serial monolayer with respect to their morphological and biosynthetic phenotype. They change from a round to a flattened fibroblast-like shape, and ***collagen***

I is secreted instead of the cartilage-specific ***collagen*** II. was analysed in detail the time course of dedifferentiation of mature bovine articular ***chondrocytes*** in monolayer for up to 32 weeks. ***chondrocytes*** AB dedifferentiate in serial monolayer with Assessment of RNA expression by reverse transcription-PCR led to the identification of two novel phenotypical markers, the ***cartilage*** ***oligomeric*** ***matrix*** ***protein*** (COMP) and ***oligomeric***

collagen

IX, which are down-regulated faster than the widely accepted marker, ***collagen***

II. The different kinetics of COMP and ***collagen***

expression suggest differential regulation at the level of transcription. Immunostaining and metabolic labelling experiments confirmed the switch in the ***collagen***

expression pattern and the rapid down-regulation of de novo synthesis of COMP and ***collagen*** IX. Culture of ***chondrocytes*** in a three-dimensional matrix is known to stabilize the chondrocytic phenotype. We maintained cells for up to 28 weeks in an alginate bead system, which prevented dedifferentiation and led to a stabilization of ***collagen*** and COMP expression. Immunohistochemical analysis of the alginate beads revealed a similar distribution of matrix proteins to that found in vivo.

Chondrocytes were transferred after a variable length of monolayer culture into the alginate matrix and the potential for redifferentiation was investigated. The re-expression of COMP and ***collagen*** IX was differentially regulated. The expression of COMP was re-induced within days after transfer into the three-dimensional matrix, while the expression of ***collagen*** IX was irreversibly down-regulated. summary, these results demonstrate that the potential for redifferentiation decreases with increasing length of monolayer culture and show that the alginate bead system represents an attractive in vitro model to study the ***chondrocyte*** de- and re-differentiation processes, as well as extracellular matrix assembly. L23 ANSWER 8 OF 12 MEDLINE on STN 2001431622 ACCESSION NUMBER: MEDLINE DOCUMENT NUMBER: 21372013 PubMed ID: 11478845 TITLE: Chondrogenic differentiation of mesenchymal stem cells from bone marrow: differentiation-dependent gene expression of matrix components. Barry F; Boynton R E; Liu B; Murphy J M
Osiris Therapeutics, Inc., 2001 Aliceanna Street,
Baltimore, Maryland 21231, USA.. fbarry@osiristx.com **AUTHOR:** CORPORATE SOURCE: EXPERIMENTAL CELL RESEARCH, (2001 Aug 15) 268 (2) 189-200. Journal code: 0373226. ISSN: 0014-4827. SOURCE: PUB. COUNTRY: United States DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE) LANGUAGE: English FILE SEGMENT: Priority Journals ENTRY MONTH: 200109 ENTRY DATE: Entered STN: 20010924 Last Updated on STN: 20010924 Entered Medline: 20010920 Transforming growth factor (TGF)-beta-induced chondrogenesis of ***mesenchymal*** ***stem*** ***cells*** derived from the first transforming growth factor (TGF)-beta-induced chondrogenesis of the first transforming growth AB ***cells*** derived from bone marrow involves the rapid deposition of a cartilage-specific extracellular matrix. The sequential events in this pathway leading from the undifferentiated stem cell to a mature ***chondrocyte*** we

investigated by analysis of key matrix elements. Differentiation was

rapidly induced in cells cultured in the presence of TGF-beta 3 or -beta 2 and was accompanied by the early expression of fibromodulin a ***cartilage*** ***oligomeric*** ***matrix*** ***protein*** An increase in aggrecan and versican core protein synthesis defined an intermediate stage, which also involved the small leucine-rich proteoglycans decorin and biglycan. This was followed by the appearance of type II ***collagen*** and chondroadherin. The pathway was also of type II ***collagen*** and chondroadherin. The pathway was also characterized by the appearance of type X ***collagen***, usually associated with hypertrophic cartilage. There was also a change in the pattern of sulfation of chondroitin sulfate, with a progressive increase in the proportion of 6-sulfated species. The major proportion of newly synthesized glycosaminoglycan was part of an aggregating proteoglycan network. These data allow us to define the phenotype of the differentiated cell and to understand in greater detail the sequential process of matrix assembly. Copyright 2001 Academic Press.

L23 ANSWER 9 OF 12 MEDLINE on STN 2001349628 ACCESSION NUMBER: MEDLINE

21305865 DOCUMENT NUMBER: PubMed ID: 11412822

TITLE: Cartilage and bone biological markers in the synovial fluid

of osteoarthritic patients after hyaluronan injections in

AUTHOR:

Herrero-Beaumont G; Guerrero R; Sanchez-Pernaute O; Acebes C; Palacios I; Mas S; Rodriguez I; Egido J; Vivanco F Inflammation Research Unit, Fundacion Jimenez Diaz, Avda. CORPORATE SOURCE:

Reyes Catolicos 2, 28040 Madrid, Spain.. gherrero@fjd.es CLINICA CHIMICA ACTA, (2001 Jun) 308 (1-2) 107-15. Journal code: 1302422. ISSN: 0009-8981. SOURCE:

PUB. COUNTRY: Netherlands DOCUMENT TYPE: (CLINICAL TRIAL)

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

Priority Journals FILE SEGMENT:

ENTRY MONTH: 200107

ENTRY DATE: Entered STN: 20010730

Last Updated on STN: 20010730

Entered Medline: 20010726
OBJECTIVE: To evaluate synovial fluid levels of cartilage and bone biological markers after repetitive intra-articular injections of sodium hyaluronate (HA) in knee osteoarthritis (OA) patients. METHODS: Twenty patients with knee OA were evaluated before and after 5 weekly injections of HA. To study cartilage and bone biological markers, synovial fluid and urine samples were collected simultaneously with the first (FI=week O) and before the last injection (LI=week 4) of HA. Not commercially available markers (***cartilage*** ***oligomeric*** ***matrix***

protein (COMP), proteoglycan monomers and cyanogen bromide peptide 11 of the type II ***collagen*** chains (alpha (II) CA11B)) were determined by an indirect inhibition ELISA developed and standardized in our laboratory. RESULTS: We found a significant reduction in levels of proteoglycan monomers (30+/-16 vs. 22+/-10 microg/ml, p<0.05), an increase in COMP concentration (2.9+/-0.9 vs. 3.6+/-0.9 microg/ml, p<0.05) and osteocalcin (BGP) levels (8.7+/-8 vs. 11.9+/-9 ng/ml, p<0.05). No significant changes were observed in the levels of alpha (II)CB11B) metalloproteinase-1 (MMP-1) or pyridinium cross-link/creatinine (Pyr/Cr). CONCLUSIONS: HA could elicit an indirect response on the cartilage and bone metabolism due to the increased overuse of the joint caused by the analgesic effect of HA. However, a direct HA action on the metabolism of ***chondrocytes*** must not be ruled out.

L23 ANSWER 10 OF 12 MEDLINE on STN ACCESSION NUMBER: 2000122597 MEDLINE

DOCUMENT NUMBER: 20122597 PubMed ID: 10655510

TITLE: A mutation in the alpha 3 chain of type IX collagen causes

autosomal dominant multiple epiphyseal dysplasia with mild

AUTHOR:

Bonnemann C G; Cox G F; Shapiro F; Wu J J; Feener C A; Thompson T G; Anthony D C; Eyre D R; Darras B T; Kunkel L M Department of Medicine (Genetics), Children's Hospital,

CORPORATE SOURCE:

Boston, MA 02115, USA.

P30-HD18655 (NICHD) CONTRACT NUMBER:

SOURCE: PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES OF THE

UNITED STATES OF AMERICA, (2000 Feb 1) 97 (3) 1212-7. Journal code: 7505876. ISSN: 0027-8424.

PUB. COUNTRY: **United States**

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: ENTRY MONTH: **ENTRY DATE:**

Priority Journals 200003

Entered STN: 20000314

Last Updated on STN: 20000314
Entered Medline: 20000302
Multiple epiphyseal dysplasia (MED) is a degenerative cartilage condition AB shown in some cases to be caused by mutations in genes encoding ***cartilage*** ***oligomeric*** ***matrix*** ***

protein ***collagen*** We studied a family with autosomal dominant MED affecting predominantly the knee joints and a mild proximal myopathy. Genetic linkage to the COL9A3 locus on chromosome 20q13.3 was established with a peak log(10) odds ratio for linkage score of 3.87 for markers D20S93 and D20S164. Reverse transcription-PCR performed on the muscle biopsy revealed aberrant mRNA lacking exon 3, which predicted a protein lacking 12 amino acids from the COL3 domain of alpha3(IX)

collagen Direct sequencing of genomic DNA confirmed the presentations of the collagen of the

collagen . Direct sequencing of genomic DNA confirmed the presence of a splice acceptor mutation in intron 2 of the COL9A3 gene (intervening sequence 2, G-A, -1) only in affected family members. By electron microscopy, ***chondrocytes*** from epiphyseal cartilage exhib from epiphyseal cartilage exhibited dilated rough endoplasmic reticulum containing linear lamellae of alternating electron-dense and electron-lucent material, reflecting abnormal processing of mutant protein. Type IX ***collagen*** chains appeared normal in size and quantity but showed defective cross-linking by Western blotting. The novel phenotype of MED and mild myopathy is likely caused by a dominant-negative effect of the exon 3-skipping mutation in the COLONG data. the COL9A3 gene. Patients with MED and a waddling gait but minimal radiographic hip involvement should be evaluated for a primary myopathy

L23 ANSWER 11 OF 12 MEDLINE on STN 1998049569 ACCESSION NUMBER: MEDLINE

DOCUMENT NUMBER: 98049569 PubMed ID: 9388247

and a mutation in type IX

The fate of cartilage oligomeric matrix protein is TITLE:

determined by the cell type in the case of a novel mutation

in pseudoachondroplasia.

AUTHOR: Maddox B K; Keene D R; Sakai L Y; Charbonneau N L; Morris N

collagen

P; Ridgway C C; Boswell B A; Sussman M D; Horton W A; Bachinger H P; Hecht J T

Research Department, Shriners Hospital for Children, Portland, Oregon 97201, USA. CORPORATE SOURCE:

SOURCE: JOURNAL OF BIOLOGICAL CHEMISTRY, (1997 Dec 5) 272 (49)

30993-7.

Journal code: 2985121R. ISSN: 0021-9258.

PUB. COUNTRY: **United States**

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: ENTRY MONTH: Priority Journals

199801

ENTRY DATE: Entered STN: 19980122

Last Updated on STN: 19990129 Entered Medline: 19980108

We have identified a novel missense mutation in a pseudoachondroplasia (PSACH) patient in one of the type III repeats of ***cartilage***

oligomeric ***matrix*** ***protein*** (COMP). Enlarge AB lamellar rough endoplasmic reticulum vesicles were shown to contain accumulated COMP along with type IX ***collagen***, a cartilage-specific component. COMP was secreted and assembled normally into the extracellular matrix of tendon, demonstrating that the accumulation of COMP in ***chondrocytes*** was a cell-specific phenomenon. We believe that the intracellular storage of COMP causes a nonspecific aggregation of cartilage-specific molecules and results in a cartilage matrix deficient in required structural components leading to impaired cartilage growth and maintenance. These data support a common pathogenetic mechanism behind two clinically related chondrodysplasias, PSACH and multiple epiphyseal dysplasia.

ANSWER 12 OF 12 CAPLUS COPYRIGHT 2003 ACS ON STN

ACCESSION NUMBER:

2002:933360 CAPLUS

DOCUMENT NUMBER:

138:382725

TITLE: Effects of overexpression of membrane-bound

transferrin-like protein (MTf) on chondrogenic

differentiation in vitro

AUTHOR(S):

Suardita, Ketut; Fujimoto, Katsumi; Oda, Ryo; Shimazu, Atsushi; Miyazaki, Kazuko; Kawamoto, Takeshi; Kato,

Graduate School of Biomedical Sciences, Departments of CORPORATE SOURCE: Dental and Medical Biochemistry, Hiroshima University,

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Minami-ku Hiroshima, 734-8553, Japan Journal Biological Chemistry (2002)
                                                Biological Chemistry (2002), 2
                                                                                             50),
 SOURCE:
                                  48579-48586
                                  CODEN: JBCHA3; ISSN: 0021-9258
 PUBLISHER:
                                  American Society for Biochemistry and Molecular
                                  Biology
 DOCUMENT TYPE:
                                  Journal
 LANGUAGE:
                                  English
       Membrane-bound transferrin-like protein (MTf) is expressed in parallel
       with the expression of cartilage-characteristic genes during
       differentiation of chondrocytes, and the MTf level is much higher in
       cartilage than in other tissues. To investigate the role of MTf in cartilage, we examd the effects of growth factors on MTf expression in mouse prechondrogenic ATDC5 cells and the effect of MTf overexpression on
       differentiation of ATDC5 and mouse pluripotent mesenchymal C3H1OT1/2
                  In ATDC5 cultures, bone morphogenetic protein-2 and transforming
       growth factor-.beta. as well as insulin induced MTf mRNA expression when
       these peptides induced chondrogenic differentiation. Forced expression of
       rabbit MTf in ATDC5 cells induced aggrecan, type II collagen, matrilin-1, type X collagen mRNAs, and cell-shape changes from fibroblastic cells to
       spherical chondrocytes. Accordingly, the synthesis and accumulation of
       proteoglycans were higher in MTf-expressing cultures than in control cultures. These effects of MTf overexpression correlated with the MTf protein level on the cell surface and decreased in the presence of
       anti-MTf antibody. However, the aggrecan mRNA level in the ATDC5 cells overexpressing MTf was lower than that in Wild type ATDC5 cells exposed to
       10 .mu.g/mL insulin. MTf overexpression in C3H1OT1/2 cells also induced
       aggrecan and/or type II collagen mRNA but not the spherical phenotype. These findings suggest that the expression of MTf on the cell surface facilitates the differentiation of prechondrogenic cells, although MTf overexpression alone seems to be insufficient to commit pluripotent
       mesenchymal cells to the chondrocyte lineage.
                                          THERE ARE 38 CITED REFERENCES AVAILABLE FOR THIS
REFERENCE COUNT:
                                          RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT
=> d his
       (FILE 'HOME' ENTERED AT 10:00:47 ON 13 OCT 2003)
       FILE 'MEDLINE, CAPLUS, BIOSIS, EMBASE, SCISEARCH, AGRICOLA' ENTERED AT
       10:01:08 ON 13 OCT 2003
L1
               1106 S HCOMP OR (CARTILAGE OLIGOMERIC MATRIX PROTEIN) OR (THROMBOSPO
L2
L3
              55314 S TRYPSIN (P) (CLEAV? OR DIGEST?)
35722 S (50 KDA) OR (55 KDA) OR (62 KDA) OR (67 KDA)
L4
                   0 S L2 (P) L3 (P) L1
            269998 S ELISA
L5
L6
               8257 S L5 (P) KIT
                   3 S L1 (P) L6
L7
L8
                   1 DUPLICATE REMOVE L7 (2 DUPLICATES REMOVED)
L9
                292 S L1 (P) HUMAN
                   0 S L9 (P) L6
L10
L11
            515579 S (BIOLOGICAL MATRIX) OR (TREATED CARTILAGE) OR (BONE MATRIX) O
                249 S L1 (P) L11
                 11 S L12 (P) COMPOSITION
4 DUPLICATE REMOVE L13 (7 DUPLICATES REMOVED)
L14
L15
                 10 S L12 (P) (PURIFIED)
                   2 DUPLICATE REMOVE L15 (8 DUPLICATES REMOVED)
L17
                     S L16 NOT L14
L18
             51010 S CHONDROCYTE OR (MESENCHYMAL STEM CELL)
L19
                831 S DIFFERENTIATION AGENT
L20
            210357 S (VITAMIN D) OR (RETINOIC ACID)
L21
                 78 DUPLICATE REMOVE L12 (171 DUPLICATES REMOVED)
L22
                 14 S L21 (P) L18
                 12 S L22 NOT (L17 OR L14)
L23
=> s 121 (p) (119 or 120)
PROXIMITY OPERATOR LEVEL NOT CONSISTENT WITH
FIELD CODE - 'AND' OPERATOR ASSUMED 'L168 (P)
PROXIMITY OPERATOR LEVEL NOT CONSISTENT WITH FIELD CODE - 'AND' OPERATOR ASSUMED 'L172 (P)
                 0 L21 (P) (L19 OR L20)
=> d his
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10:01:08 ON 13 OCT 2003
             1106 S HCOMP OR (CARTILAGE OLIGOMERIC MATRIX PROTEIN) OR (THROMBOSPO 55314 S TRYPSIN (P) (CLEAV? OR DIGEST?) 35722 S (50 KDA) OR (55 KDA) OR (62 KDA) OR (67 KDA)
 L1
 L3
                 0 S L2 (P) L3 (P) L1
 L4
 L5
            269998 S ELISA
 L6
              8257 S L5 (P) KIT
 L7
                 3 S L1 (P) L6
                 1 DUPLICATE REMOVE L7 (2 DUPLICATES REMOVED)
 L8
 L9
               292 S L1 (P) HUMAN
                 0 S L9 (P) L6
 L10
           515579 S (BIOLOGICAL MATRIX) OR (TREATED CARTILAGE) OR (BONE MATRIX) O
 L11
               249 S L1 (P) L11
11 S L12 (P) COMPOSITION
 L12
 L13
                 4 DUPLICATE REMOVE L13 (7 DUPLICATES REMOVED)
 L14
                10 S L12 (P) (PURIFIED)
 L15
 L16
                 2 DUPLICATE REMOVE L15 (8 DUPLICATES REMOVED)
                 2 S L16 NOT L14
 L17
             51010 S CHONDROCYTE OR (MESENCHYMAL STEM CELL)
 L18
 L19
               831 S DIFFERENTIATION AGENT
           210357 S (VITAMIN D) OR (RETINOIC ACID)
                78 DUPLICATE REMOVE L12 (171 DUPLICATES REMOVED)
 L22
                14 S L21 (P) L18
                12 S L22 NOT (L17 OR L14)
 L23
<u>- L24</u>
                 0 S L21 (P) (L19 OR L20)
 => log y
-COST-IN-U-S. DOLLARS
                                                          SINCE FILE
                                                                            TOTAL
                                                               ENTRY
                                                                          SESSION
 FULL ESTIMATED COST
                                                               108.81
                                                                           109.02
 DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)
                                                          SINCE FILE
                                                                            TOTAL
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ENTRY

-0.65

SESSION

-0.65

FILE 'MEDLINE, CAPLUS, BIOSIS EMBASE, SCISEARCH, AGRICOLA' ENTERED AT

STN INTERNATIONAL LOGOFF AT 10:17:31 ON 13 OCT 2003

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FILE 'MEDLINE' ENTERED AT 10:31:5 N 13 OCT 2003
 FILE 'CAPLUS' ENTERED AT 10:31:56 ON 13 OCT 2003
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 FILE 'SCISEARCH' ENTERED AT 10:31:56 ON 13 OCT 2003
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 FILE 'AGRICOLA' ENTERED AT 10:31:56 ON 13 OCT 2003
 => s hcomp or (cartilage oligomeric matrix protein) or thromospondin-5
              1097 HCOMP OR (CARTILAGE OLIGOMERIC MATRIX PROTEIN) OR THROMOSPONDIN-
=> s chen hui/au
L2
               672 CHEN HUI/AU
=> s lawler john/au
                13 LAWLER JOHN/AU
L3
=> s 11 and (12 or 13)
                  7 L1 AND (L2 OR L3)
=> duplicate remove 14
DUPLICATE PREFERENCE IS 'CAPLUS, BIOSIS'
KEEP DUPLICATES FROM MORE THAN ONE FILE? Y/(N):n
PROCESSING COMPLETED FOR L4
                   4 DUPLICATE REMOVE L4 (3 DUPLICATES REMOVED)
=> d 15 1-4 ibib abs
       ANSWER 1 OF 4 CAPLUS COPYRIGHT 2003 ACS on STN DUPLICATE 1
ACCESSION NUMBER:
                                 2001:79527 CAPLUS
                                 134:205377
DOCUMENT NUMBER:
TITLE:
                                    ***Cartilage***
                                                              ***oligomeric***
                                    ***matrix***
                                                          ***protein***
                                                                                (thrombospondin-5) is
                                 expressed by human vascular smooth muscle cells
AUTHOR(S):
                                 Riessen, Reimer; Fenchel, Michael;
                                                                                  ***Chen, Hui***
                                 ; Axel, Dorothea I.; Karsch, Karl R.; Lawler, Jack
Department of Medicine III (Cardiology), University of
Tubingen, Tubingen, 72076, Germany
Arteriosclerosis, Thrombosis, and Vascular Biology
(2001), 21(1), 47-54
CODEN: ATVBFA; ISSN: 1079-5642
CORPORATE SOURCE:
SOURCE:
PUBLISHER:
                                 Lippincott Williams & Wilkins
DOCUMENT TYPE:
                                 Journal
                                 English
    ***oligomeric***
LANGUAGE:
         ***Cartilage***
                                                              ***matrix***
                                                                                     ***protein***
      (COMP/thrombospondin [TSP]-5) belongs to the thrombospondin gene family and is an extracellular glycoprotein found predominantly in cartilage and tendon. To date, there is limited evidence of COMP/TSP-5 expression
      outside of the skeletal system. The aim of the present study was to
       investigate the expression of COMP/TSP-5 in cultured human vascular smooth
      muscle cells and human arteries. COMP/TSP-5 mRNA and protein expression was detected in cultured human vascular smooth muscle cells with both
      Northern blotting and immunopptn. Serum, as well as transforming growth factor (TGF)-/.beta.1 and TGF-/33, stimulated COMP/TSP-5 mRNA expression. COMP/TSP-5 was detected in normal as well as atherosclerotic and
       restenotic human arteries with immunohistochem. The majority of
      COMP/TSP-5 was expressed in close proximity to vascular smooth muscle
      cells. In vitro attachment assays demonstrated strong adhesion of smooth
      muscle cells to COMP/TSP-5- coated surfaces, with the majority of cells
       spreading and forming stress fibers. In addn., COMP/TSP-5 supported the
      migration of smooth muscle cells in vitro. The present study shows that COMP/TSP-5 is present in human arteries and may play a role in the adhesion and migration of vascular smooth muscle cells during
      vasculogenesis and in vascular disease settings such as atherosclerosis.
REFERENCE COUNT:
                                 39
                                        THERE ARE 39 CITED REFERENCES AVAILABLE FOR THIS
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```
ANSWER 2 OF 4 CAPLUS COPYRIGHT 2003 ACS on STN DUPLICATE 2
 L5
                                               2000:627777 CAPLUS
 ACCESSION NUMBER:
 DOCUMENT NUMBER:
                                               133:331101
                                                  ***Cartilage***
                                                                                      ***oligomeric***
 TITLE:
                                                  ***matrix***
                                                                                 ***protein*** is a calcium-binding
                                              protein, and a mutation in its type 3 repeats causes
                                              conformational changes
                                                  ***Chen, Hui***
AUTHOR(S):
                                                                                 ; Deere, Michelle; Hecht,
                                              Jacqueline T.; Lawler, Jack
                                              Division of Tumor Biology and Angiogenesis, Department of Pathology, Beth Israel Deaconess Medical Center and
CORPORATE SOURCE:
                                              Harvard Medical School, Boston, MA, 02215, USA
Journal of Biological Chemistry (2000), 275(34),
SOURCE:
                                              26538-26544
                                              CODEN: JBCHA3; ISSN: 0021-9258
PUBLISHER:
                                              American Society for Biochemistry and Molecular
DOCUMENT TYPE:
                                              Journal
LANGUAGE:
                                              English
         Mutations in residues in the type 3 calcium-binding repeats and C-terminal globular region of ***cartilage*** ***oligomeric*** ***matrix***
         ***protein*** (COMP) lead to two skeletal dysplasias,
pseudoachondroplasia and multiple epiphyseal dysplasia. It has been
hypothesized that these mutations cause COMP to misfold and to be retained
         in the endoplasmic reticulum. However, this hypothesis is not supported by previous reports that COMP, when purified in the presence of EDTA, shows no obvious difference in electron microscopic appearance in the
         presence or absence of calcium ions. Since this discrepancy may be due to the removal of calcium during purifn., we have expressed wild-type COMP and the most common mutant form found in pseudoachondroplasia, MUT3, using a mammalian expression system and have purified both proteins in the presence of calcium. Both proteins are expressed as pentamers. Direct calcium binding expts. demonstrate that wild-type COMP, when purified in the presence of calcium, is a calcium-binding protein. Rotary shadowing electron microscopy and limited trypsin digestion at various calcium concess show that there are conformational changes associd with calcium
         concns. show that there are conformational changes assocd. with calcium
         binding to COMP. Whereas COMP exists in a more compact conformation in the presence of calcium, it shows a more extended conformation when calcium is removed. MUT3, with a single aspartic acid deletion in the type 3 repeats, binds less calcium and expenses a intermediate
         conformation between the calcium-replete and calcium-depleted forms of
         COMP. In conclusion, we show that a single mutation in the type 3 repeats
         of COMP causes the mutant protein to misfold. Our data demonstrate the
         importance of calcium binding to the structure of COMP and provide a
         plausible explanation for the observation that mutations in the type 3
         repeats and C-terminal globular region lead to pseudoachondroplasia.
ENCE COUNT: 39 THERE ARE 39 CITED REFERENCES AVAILABLE FOR THIS
REFERENCE COUNT:
                                                        RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT
         ANSWER 3 OF 4 CAPLUS COPYRIGHT 2003 ACS on STN DUPLICATE 3
                                             1998:554609 CAPLUS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                                             129:258910
TITLE:
                                                                                     ***cartilage***
                                             Characterization of
                                                 ***oligomeric***
                                                                                      ***matrix***
                                                                                                                      ***protein***
                                             (COMP) in human normal and pseudoachondroplasia musculoskeletal tissues
                                             Hecht, Jacqueline T.; Deere, Michelle; Putnam, Elizabeth; Cole, William; Vertel, Barbara; ***Chen,***
Hui***; Lawler, Jack
AUTHOR(S):
                                             Hui*** ; Lawler, Jack
Department of Pediatrics, University of Texas Medical
CORPORATE SOURCE:
                                             School at Houston, Houston, TX, USA
Matrix Biology (1998), 17(4), 269-278
CODEN: MTBOEC; ISSN: 0945-053X
Gustav Fischer Verlag
SOURCE:
PUBLISHER:
DOCUMENT TYPE:
                                             Journal
                                             English
***oligomeric***
LANGUAGE:
            ***Cartilage***
ΑB
                                                                                       ***matrix***
                                                                                                                      ***protein***
         (COMP), the fifth member of the -thrombospondin gene family, is an
        extracellular matrix calcium-binding protein. The importance of COMP is underscored by the finding that mutations in COMP cause the human dwarfing condition, pseudoachondroplasia (PSACH). Here, we report the results of human tissue distribution and cell secretion studies of human COMP. COMP is expressed and secreted by cultured monolayer chondrocyte, tendon and ligament cells, and COMP secretion is not restricted to a differentiated chondrocyte phenotyne. Whereas COMP is retained in the endonlasmic
        chondrocyte phenotype. Whereas COMP is retained in the endoplasmic
```

reticulum that accumulates within PSACH chondrocytes in vivo, COMP is not retained intracellularly in dedifferentiated PSACH chondrocytes in cultures. These results lend further support to the hypothesis that retention of COMP is related to the terminal PSACH chondrocyte phenotype, processing of proteins related to extracellular matrix formation, and maintenance in cartilage.

REFERENCE COUNT:

THERE ARE 28 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. on STN 1996:56780 BIOSIS ANSWER 4 OF 4 L5

ACCESSION NUMBER: PREV199698628915 DOCUMENT NUMBER:

TITLE:

Inhibition of angiogenesis by thrombospondin-2. Volpert, Olga V. (1); Tolsma, Sara S. (1); Pellerin, AUTHOR(S):

Sylvie; Feige, Jean-Jacques; ***Chen, Hui***; Mosher,

Deane F.; Bouck, Noel (1)

(1) Dep. Microbiol-Immunol., Univ. Med. Sch., Chicago, IL CORPORATE SOURCE:

USA

SOURCE: Biochemical and Biophysical Research Communications, (1995)

Vol. 217, No. 1, pp. 326-332. ISSN: 0006-291X.

DOCUMENT TYPE: LANGUAGE:

Article English

AB To assess the ability of proteins of the thrombospondin family to inhibit

angiogenesis, recombinant murine thrombospondin-2, bovine

thrombospondin-2/CISP and thrombospondin-5/COMP were purified and tested for ability to block the migration of capillary endothelial cells towards a variety of inducers and to inhibit neovascularization induced in the rat-cornea. Both preparations of thrombospondin-2 were active inhibitors in vitro and in vivo whereas thrombospondin-5/COMP was inactive. These results define thrombospondin-2 as a newly identified naturally occurring inhibitor of angiogenesis and suggest that the properdin-like type 1 modules that it shares with antiangiogenic thrombospondin-1 and are missing in thrombospondin-5/COMP could contribute to this activity.

=> d his

L1

L2

L3

(FILE 'HOME' ENTERED AT 10:31:29 ON 13 OCT 2003)

FILE 'MEDLINE, CAPLUS, BIOSIS, EMBASE, SCISEARCH, AGRICOLA' ENTERED AT 10:31:56 ON 13 OCT 2003

1097 S HCOMP OR (CARTILAGE OLIGOMERIC MATRIX PROTEIN) OR THROMOSPOND

672 S CHEN HUI/AU

13 S LAWLER JOHN/AU S L1 AND (L2 OR L3)

4 DUPLICATE REMOVE L4 (3 DUPLICATES REMOVED)

=> log y

COST IN U.S. DOLLARS SINCE FILE TOTAL **ENTRY SESSION** FULL ESTIMATED COST 29.64 29.85

DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS) SINCE FILE TOTAL **ENTRY SESSION** CA SUBSCRIBER PRICE -1.95-1.95

STN INTERNATIONAL LOGOFF AT 10:34:17 ON 13 OCT 2003